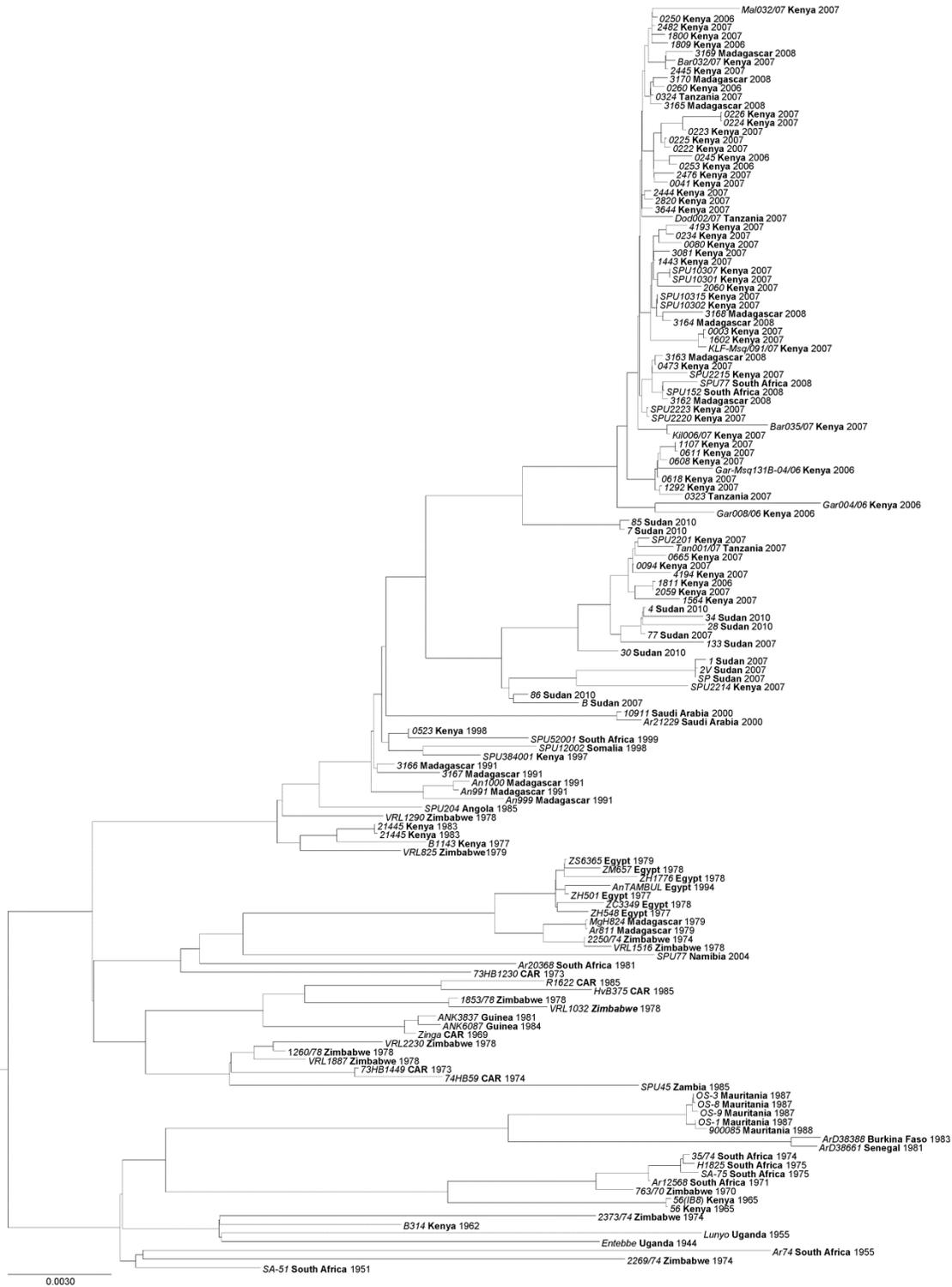


Rift Valley Fever, Sudan, 2007 and 2010

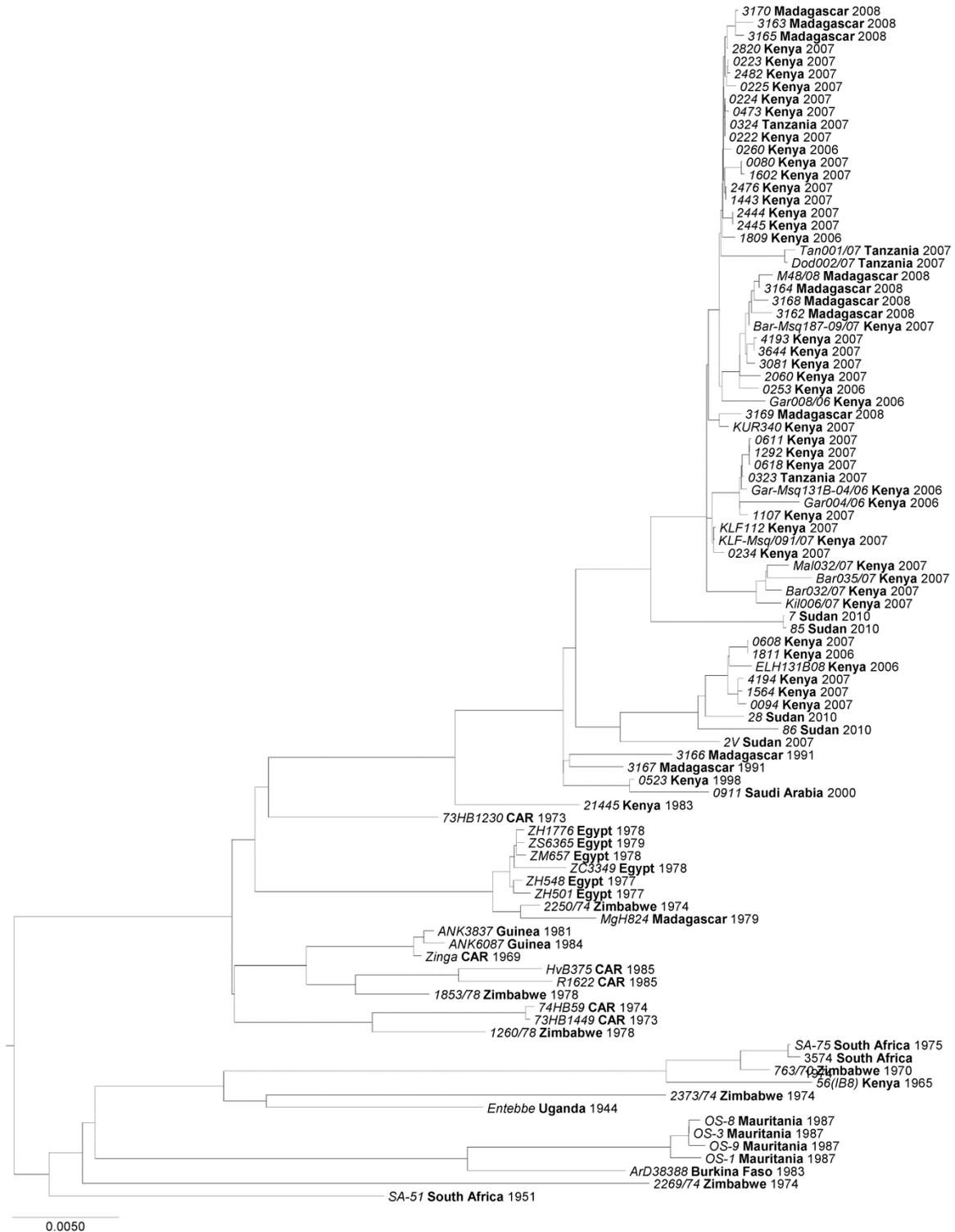
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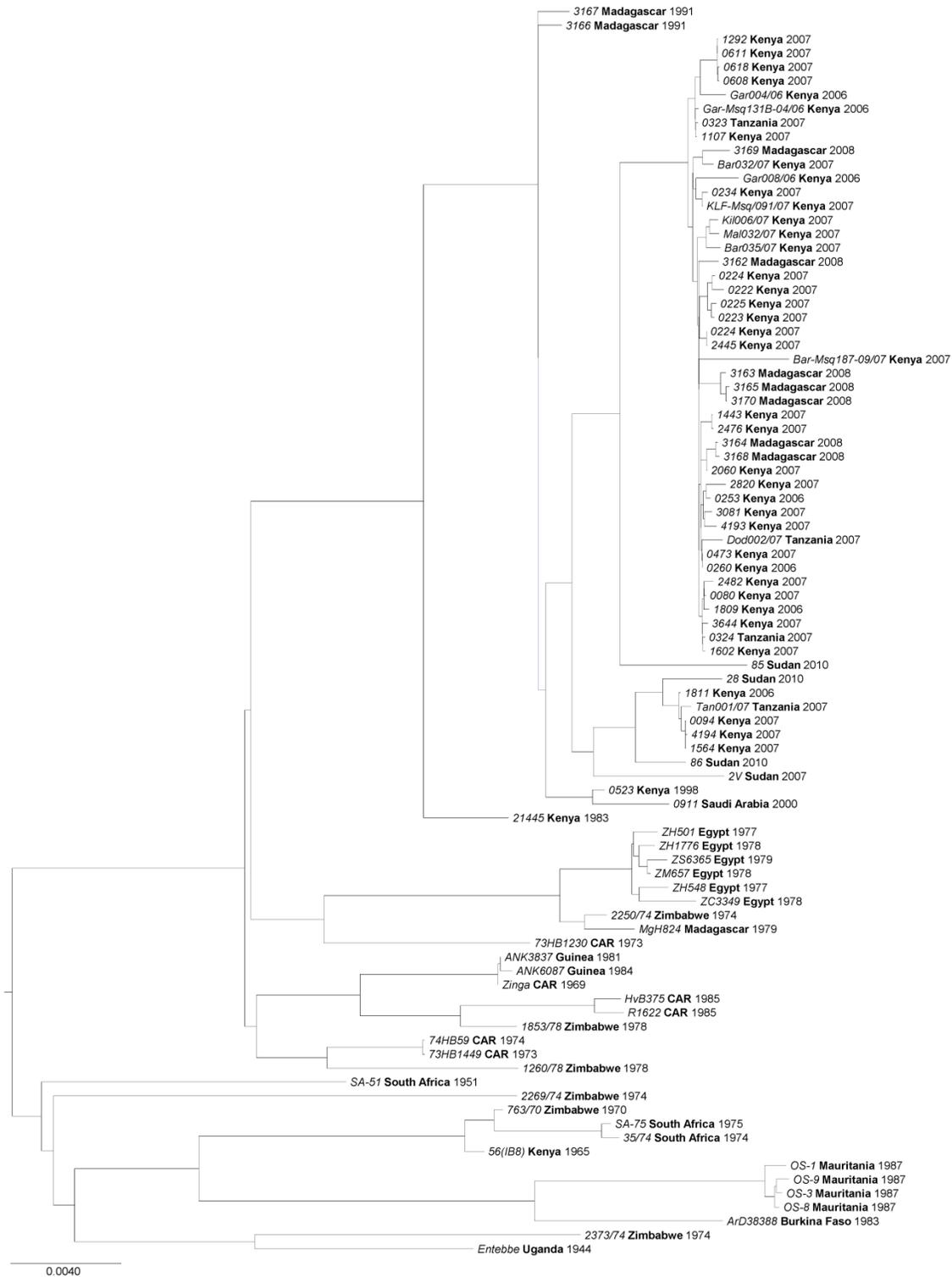
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Technical Appendix Figure 1. Phylogenetic analysis of complete Rift Valley fever virus S (small) segment sequences represented as a maximum clade credibility tree. Scale bar represents substitutions per site per year.



Technical Appendix Figure 2. Phylogenetic analysis of complete Rift Valley fever virus M (medium) segment sequences represented as a maximum clade credibility tree. Scale bar represents substitutions per site per year.



Technical Appendix Figure 3. Phylogenetic analysis of complete Rift Valley fever virus L (large) segment sequences represented as a maximum clade credibility tree. Scale bar represents substitutions per site per year.